
Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: [year=2010; month=5; day=26; hr=13; min=22; sec=8; ms=534;]

Validated By CRFValidator v 1.0.3

Application No: 10574182 Version No: 1.0

Input Set:

Output Set:

Started: 2010-05-20 19:07:46.112

Finished: 2010-05-20 19:07:48.523

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 411 ms

Total Warnings: 26

Total Errors: 0

No. of SeqIDs Defined: 26

Actual SeqID Count: 26

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W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(3)
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W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(5)
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W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(8)
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W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(10)
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W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(19)
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Input Set:

Output Set:

Started: 2010-05-20 19:07:46.112 **Finished:** 2010-05-20 19:07:48.523

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 411 ms

Total Warnings: 26

Total Errors: 0

No. of SeqIDs Defined: 26

Actual SeqID Count: 26

Error code Error Description

This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

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                            40
                                                45
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<212> PRT

<213> homo sapien

<400> 4

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Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro

2.0 2.5

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<400> 8

Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val 5 10 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro 25 20 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu 40 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu 55 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly 65 70 75 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly 85 90 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn 100 105 Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln 120 Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu 135 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His 145 150 155 Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg 165 170 175 Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu 180 185 190 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr 200 His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser His Ser Arg 215 Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg 225 230 235 240 Gln Leu

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Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
                           40
                                                45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
                        55
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
                   7.0
                                       7.5
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
               8.5
                                    90
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
                                105
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
                            120
                                                125
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
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                                           140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
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                                       155
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
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Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu Ser Glu
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His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser His Ser Arg
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<212> PRT

<213> homo sapien

<400> 12

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Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro 20 25 30

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Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu 50 55 60

Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly 65 70 75 80

Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
85 90 95

Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn 100 105 110

Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln 115 120 125

Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu